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## Recra LabNet Philadelphia Analytical Report

**Client :** TNU-HANFORD B98-063

**RFW# :** 9808L205

**SDG/SAF:** H0186/B98-063

**W.O.# :** 10985-001-001-9999-00

**Date Received :** 08-03-98

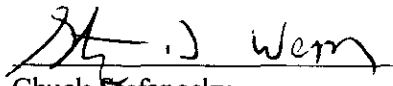
### PCB

One (1) water sample was collected on 08-03-98.

The sample and its associated QC samples were extracted on 08-07-98 and analyzed on 08-17-98 according to Recra OPs based on SW846, 3rd Edition, Method 3520 and Method 8081.

The following is a summary of the QC results accompanying the sample results and a description of any problems encountered during their analyses:

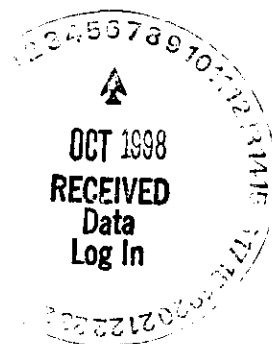
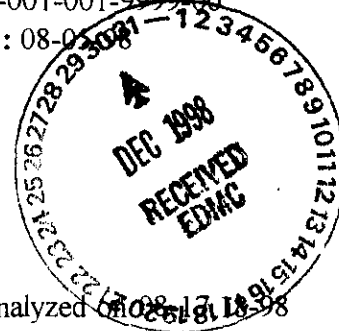
1. The cooler temperature upon receipt has been recorded on the chain-of-custody.
2. The required holding time for extraction was not met because samples were received in the Lionville laboratory outside of extraction holding time; however, all analysis holding time were met.
3. The sample and its associated QC samples received a sulfuric acid and sulfur cleanup.
4. The method blank was below the reporting limits for all target compounds.
5. All surrogate recoveries were within acceptance criteria.
6. The blank spike recovery was within acceptance criteria.
7. All matrix spike recoveries were within acceptance criteria.
8. All initial calibrations associated with this data set were within acceptance criteria.
9. All continuing calibration standards analyzed prior to sample extracts were within acceptance criteria.

  
Chuck Stefanosky  
Laboratory Director  
Lionville Analytical Laboratory

soinpcb\word6\0 pest-pcb\tnu8p205.doc

The results presented in this report relate only to the analytical testing and conditions of the samples at receipt and during storage. All pages of this report are integral parts of the analytical data. Therefore, this report should only be reproduced in its entirety of 7 pages.

08-03-98  
Date



## GLOSSARY OF PESTICIDE/PCB DATA

### DATA QUALIFIERS

- U** = Indicates that the compound was analyzed for but not detected. The minimum detection limit for the sample (not the method detection limit) is reported with the U (e.g., 10U).
- J** = Indicates an estimated value. This flag is used in cases where a target analyte is detected at a level less than the lower quantification level. If the limit of quantification is 10 ug/L and a concentration of 3 ug/L is calculated, it is reported as 3J.
- B** = This flag is used when the analyte is found in the associated blank as well as in the sample. It indicates possible/probable blank contamination.
- E** = Indicates that the compound was detected beyond the calibration range and was subsequently analyzed at a dilution.
- I** = Interference.

### ABBREVIATIONS

- BS** = Indicates blank spike in which reagent grade water is spiked with the CLP matrix spiking solutions and carried through all the steps in the method. Spike recoveries are reported.
- BSD** = Indicates blank spike duplicate.
- MS** = Indicates matrix spike.
- MSD** = Indicates matrix spike duplicate.
- DL** = Indicates that recoveries were not obtained because the extract had to be diluted for analysis.
- NA** = Not Applicable.
- DF** = Dilution Factor.
- NR** = Not Required.
- SP** = Indicates Spiked Compound.

## GLOSSARY OF PESTICIDE/PCB DATA

- P**     =     This flag is used for a pesticide/Aroclor target analyte when there is greater than 25% difference for detected concentrations between the two GC columns (see Form X). The lower of the two values is reported on Form I and flagged with a "P".
- D**     =     This flag identifies all compounds identified in an analysis at a secondary dilution factor.
- C**     =     This flag applies to a compound that has been confirmed by GC/MS.

Recra LabNet - Lionville Laboratory

PCBs by GC

Report Date: 08/18/98 11:41

RFW Batch Number: 9808L205

Client: TNU-HANFORD B98-0063

Work Order: 10985001001 Page: 1

Cust ID:		BONCD1		BONCD1		BONCD1		PBLKMI		PBLKMI BS	
Sample Information	RFW#:	001		001 MS		001 MSD		98LE1322-MB1		98LE1322-MB1	
	Matrix:	WATER		WATER		WATER		WATER		WATER	
	D.F.:	1.00		1.00		1.00		1.00		1.00	
	Units:	UG/L		UG/L		UG/L		UG/L		UG/L	
Surrogate:	Tetrachloro-m-xylene	62	%	52	%	52	%	40	%	55	%
	Decachlorobiphenyl	76	%	93	%	84	%	86	%	103	%
=====fl=====											

U= Analyzed, not detected. J= Present below detection limit. B= Present in blank. NR= Not reported. NS= Not spiked.  
 %= Percent recovery. D= Diluted out. I= Interference. NA= Not Applicable. \*= Outside of EPA CLP QC

DR  
APR 98

004

Recra LabNet - Lionville Laboratory  
PCB ANALYTICAL DATA PACKAGE FOR  
TNU-HANFORD B98-0063

DATE RECEIVED: 08/05/98

RFW LOT # :9808L205

CLIENT ID	RFW #	MTX	PREP #	COLLECTION	EXTR/PREP	ANALYSIS
BONCD1	001	W	98LE1322	08/03/98	08/07/98	08/18/98
BONCD1	001 MS	W	98LE1322	08/03/98	08/07/98	08/18/98
BONCD1	001 MSD	W	98LE1322	08/03/98	08/07/98	08/18/98

LAB QC:

PBLKMI	MB1	W	98LE1322	N/A	08/07/98	08/17/98
PBLKMI	MB1 BS	W	98LE1322	N/A	08/07/98	08/17/98

DE  
9/2/98



**RECRA  
LabNet**

Discrepancies Between  
Samples Labels and  
COC Requirements  
NOTES

Crab not found

Collector Lahlberg Nielson		Company Contact Steve Marske		Telephone No. 373-4316		Project Coordinator WEISS, R.L.		<b>Data Turnaround</b>  <div style="font-size: 2em; font-weight: bold;">21 Days</div>			
Project Designation 105 C Phase I & Phase II - Water Samples		Sampling Location 105-C		SAF No. H98-063		Method of Shipment Federal Express					
Ice Chest No.		Field Logbook No. 11-1309-1		Offsite Property No.		Bill of Lading/Air Bill No.		<div style="font-size: 3em; font-weight: bold;">100</div>			
Shipped To <i>RUN REIRA LabNet</i>											
Waste Designation						COA					

POSSIBLE SAMPLE HAZARDS/REMARKS  4235 7951 5429 <i>g.x</i>	Preservation	None	HNO <sub>3</sub> to pH 2	Cool IC	HNO <sub>3</sub> to pH 2	HNO <sub>3</sub> to pH 2	HNO <sub>3</sub> to pH 2	Cool IC	HNO <sub>3</sub> to pH 2	HNO <sub>3</sub> to pH 2	HNO <sub>3</sub> to pH 2
	Type of Container	P	aG	P	P	P	P	aG	P	P	P
	No. of Container(s)	1	1	1	1	1	1	2	2	2	3
	Volume	20ml	500ml	500ml	500ml	1000ml	1000ml	1000ml	1000ml	1000ml	500ml
Special Handling and/or Storage Cool IC				Activity Scan	Mercury 7429-91-4 (CV)	Chromium Hex 7440-47-3	ICP Metals 6010-95-9 (Supertrace) (Lead)	See item (1) on Special Instructions	See item (2) on Special Instructions	Isotopic Plutonium Isotopic Plutonium Americium 241	Selenium 74

SAMPLE ANALYSIS			
Sample No	Matrix *	Sample Date	Sample Time
BONCD1	Water	8-3-98	1610

CHAIN OF POSSESSION	Sign/Print Names		SPECIAL INSTRUCTIONS	Matrix *
Relinquished By	Date Time	Received By	(1) Gamma Spectroscopy (Cesium-137, Cobalt-60, Europium-152, Europium-154, Europium-155) (2) Gamma Spectroscopy (Water) (Cesium-137, Cobalt-60, Europium-152, Europium-154, Europium-155)  <i>Do not run Chromium-Hex 7440 unless directed by EEC sample management.</i>	S
Relinquished By	Date Time	Received By		SO
Relinquished By	Date Time	Received By		SL
Relinquished By	Date Time	Received By		W
Relinquished By	Date Time	Received By	OL	DL
Relinquished By	Date Time	Received By		AI
Relinquished By	Date Time	Received By		DS
Relinquished By	Date Time	Received By		OL
Relinquished By	Date Time	Received By		L
Relinquished By	Date Time	Received By		WL
Relinquished By	Date Time	Received By		L
Relinquished By	Date Time	Received By		AL
Relinquished By	Date Time	Received By		AL

LABORATORY SECTION	Received By	Title	Date Time
FINAL SAMPLE DISPOSITION	Disposal Method	Disposed By	Date Time

Thermo Nutech  
W.O. No. N8-08-012-7486, SDG H0186

Bechtel Hanford Inc.  
P.O. TRB-SBB-207925

## Case Narrative

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### 1.0 GENERAL

Thermo Nutech Sample Delivery Group H0186 is comprised of a single water sample designated under SAF No. B98-063 with a Project Designation of : 105-C Phase I & II - Water Samples.

The sample was received as stated on the Chain-of-Custody documents.

### 2.0 ANALYSIS NOTES

#### 2.1 Total Strontium Analyses

No problems were encountered with the analyses. All sample MDA's were less than the RDL.

#### 2.2 Isotopic Uranium Analyses

No problems were encountered with the analyses. All sample MDA's were less than the RDL.

#### 2.3 Isotopic Plutonium Analyses

No problems were encountered with the analyses. All sample MDA's were less than the RDL.

#### 2.4 Americium-241 Analyses

No problems were encountered with the analyses. All sample MDA's were less than the RDL.

#### 2.5 Gamma Scan Analyses

No problems were encountered with the analyses. All sample MDA's were less than the RDL.



**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**REPORT GUIDE**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**ABOUT THE DATA SUMMARY SECTION**

The Data Summary Section of a Data Package has all data, in several useful orders, necessary for first level, routine review of the data package for a Sample Delivery Group (SDG). This section follows the Data Package Narrative, which has an overview of the data package and a discussion of special problems. It is followed by the Raw Data Section, which has full details.

The Data Summary Section has several groups of reports:

**SAMPLE SUMMARIES**

The Sample and QC Summary Reports show all samples, including QC samples, reported in one SDG. These reports cross-reference client and lab sample identifiers.

**PREPARATION BATCH SUMMARY**

The Preparation Batch Summary Report shows all preparation batches (lab groupings reflecting how work was organized) relevant to the reported SDG with information necessary to check the completeness and consistency of the SDG.

**WORK SUMMARY**

The Work Summary Report shows all samples and work done on them relevant to the reported SDG.

**METHOD BLANKS**

The Method Blank Reports, one for each Method Blank relevant to the SDG, show all results and primary supporting information for the blanks.

**LAB CONTROL SAMPLES**

The Lab Control Sample Reports, one for each Lab Control Sample relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

**REPORT GUIDES**

Page 1

**SUMMARY DATA SECTION**

Page 1

Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98

**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**GUIDE, cont.**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**ABOUT THE DATA SUMMARY SECTION**

**DUPLICATES**

The Duplicate Reports, one for each Duplicate and Original sample pair relevant to the SDG, show all results, differences and primary supporting information for these QC samples.

**MATRIX SPIKES**

The Matrix Spike Reports, one for each Spiked and Original sample pair relevant to the SDG, show all results, recoveries and primary supporting information for these QC samples.

**DATA SHEETS**

The Data Sheet Reports, one for each client sample in the SDG, show all results and primary supporting information for these samples.

**METHOD SUMMARIES**

The Method Summary Reports, one for each test used in the SDG, show all results, QC and method performance data for one analyte on one or two pages. (A test is a short code for the method used to do certain work to the client's specification.)

**REPORT GUIDES**

The Report Guides, one for each of the above groups of reports, have documentation on how to read the associated reports.

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

Page 2

Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98

**TMA/RICHMOND**

SAMPLE DELIVERY GROUP H0186

SDG 7486Contact N. Joseph Verville**SAMPLE SUMMARY**Client HanfordContract TRB-SBB-207925Case no SDG H0186

CLIENT SAMPLE ID	LOCATION	MATRIX	LEVEL	LAB		CHAIN OF	
				SAMPLE ID	SAF NO	CUSTODY	COLLECTED
B0NCD1	105-C	LIQUID		N808012-01	B98-063	B98-063-05	08/03/98 16:10
Method Blank		LIQUID		N808012-03	B98-063		
Lab Control Sample		LIQUID		N808012-02	B98-063		
Duplicate (N808012-01)	105-C	LIQUID		N808012-04	B98-063		08/03/98 16:10

**SAMPLE SUMMARY**

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**SUMMARY DATA SECTION**

Page 3

Lab id TMANCProtocol HanfordVersion Ver 1.0Form DVD-CSVersion 3.06Report date 08/27/98

# TMA/RICHMOND

SAMPLE DELIVERY GROUP H0186

## QC SUMMARY

SDG 7486

Contact N. Joseph Verville

Client Hanford

Contract TRB-SBB-207925

Case no SDG H0186

QC BATCH	CHAIN OF CUSTODY	CLIENT SAMPLE ID	MATRIX	SOLIDS	SAMPLE AMOUNT	BASIS AMOUNT	DAYS SINCE RECEIVED	LAB COLL SAMPLE ID	DEPARTMENT SAMPLE ID
7486	B98-063-05	B0NCD1	LIQUID				08/05/98 2	N808012-01	7486-001
		Method Blank	LIQUID					N808012-03	7486-003
		Lab Control Sample	LIQUID					N808012-02	7486-002
		Duplicate (N808012-01)	LIQUID				08/05/98 2	N808012-04	7486-004

QC SUMMARY

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SUMMARY DATA SECTION

Page 4

Lab id TMANC

Protocol Hanford

Version Ver 1.0

Form DVD-QS

Version 3.06

Report date 08/27/98

## TMA/RICHMOND

SAMPLE DELIVERY GROUP H0186

SDG 7486

Contact N. Joseph Verville

## PREP BATCH SUMMARY

Client HanfordContract TRB-SBB-207925Case no SDG H0186

TEST	MATRIX	METHOD	PREPARATION	ERROR	PLANCHETS ANALYZED				QUALI-			
			BATCH	2σ %	CLIENT	MORE	RE	BLANK	LCS	DUP/ORIG	MS/ORIG	FIERS
Alpha Spectroscopy												
AM	LIQUID	Americium 241 in Water	2785-103	5.0	1			1	1	1/1		
PU	LIQUID	Plutonium, Isotopic in Water	2785-103	5.0	1			1	1	1/1		
U	LIQUID	Uranium, Isotopic in Water	2785-103	5.0	1			1	1	1/1		
Beta Counting												
SR	LIQUID	Strontium, Total in Water	2785-103	10.0	1			1	1	1/1		
Gamma Scan												
GAM	LIQUID	Gamma Emitters	2785-103	15.0	1			1	1	1/1		

Duplicates and Matrix Spikes are those with original (Client) sample in this Sample Delivery Group.

Blank and LCS planchets are those in the same preparation batch as some Client, Duplicate or Spike sample.

PREP BATCH SUMMARY

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SUMMARY DATA SECTION

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Lab id TMANCProtocol HanfordVersion Ver 1.0Form DVD-PBSVersion 3.06Report date 08/27/98

# TMA/RICHMOND

SAMPLE DELIVERY GROUP H0186

SDG 7486

Contact N. Joseph Verville

## WORK SUMMARY

Client Hanford

Contract TRB-SBB-207925

Case no SDG H0186

CLIENT SAMPLE ID		LAB SAMPLE ID									
LOCATION	MATRIX	COLLECTED		SUF-							
CUSTODY	SAF No	RECEIVED	PLANCHET	TEST	FIX	ANALYZED	REVIEWED	BY	METHOD		
B0NCD1		N808012-01	7486-001	AM		08/25/98	08/27/98	NJV	Americium 241 in Water		
105-C		08/03/98	7486-001	GAM		08/11/98	08/27/98	NJV	Gamma Emitters		
B98-063-05	B98-063	08/05/98	7486-001	PU		08/19/98	08/27/98	NJV	Plutonium, Isotopic in Water		
			7486-001	SR		08/13/98	08/27/98	NJV	Strontium, Total in Water		
			7486-001	U		08/13/98	08/27/98	NJV	Uranium, Isotopic in Water		
Method Blank		N808012-03	7486-003	AM		08/25/98	08/27/98	NJV	Americium 241 in Water		
			7486-003	GAM		08/11/98	08/27/98	NJV	Gamma Emitters		
	B98-063		7486-003	PU		08/19/98	08/27/98	NJV	Plutonium, Isotopic in Water		
			7486-003	SR		08/13/98	08/27/98	NJV	Strontium, Total in Water		
			7486-003	U		08/13/98	08/27/98	NJV	Uranium, Isotopic in Water		
Lab Control Sample		N808012-02	7486-002	AM		08/25/98	08/27/98	NJV	Americium 241 in Water		
			7486-002	GAM		08/11/98	08/27/98	NJV	Gamma Emitters		
	B98-063		7486-002	PU		08/19/98	08/27/98	NJV	Plutonium, Isotopic in Water		
			7486-002	SR		08/13/98	08/27/98	NJV	Strontium, Total in Water		
			7486-002	U		08/13/98	08/27/98	NJV	Uranium, Isotopic in Water		
Duplicate (N808012-01)		N808012-04	7486-004	AM		08/26/98	08/27/98	NJV	Americium 241 in Water		
105-C		08/03/98	7486-004	GAM		08/12/98	08/27/98	NJV	Gamma Emitters		
	B98-063	08/05/98	7486-004	PU		08/19/98	08/27/98	NJV	Plutonium, Isotopic in Water		
			7486-004	SR		08/13/98	08/27/98	NJV	Strontium, Total in Water		
			7486-004	U		08/13/98	08/27/98	NJV	Uranium, Isotopic in Water		

## COUNTS OF TESTS BY SAMPLE TYPE

TEST	SAF No	METHOD	REFERENCE	CLIENT	MORE	RE	BLANK	LCS	DUP	SPIKE	TOTAL
AM	B98-063	Americium 241 in Water	AM/CMPLATE	1			1	1	1		4
GAM	B98-063	Gamma Emitters	GAMMAHI	1			1	1	1		4
PU	B98-063	Plutonium, Isotopic in Water	PUPLATE	1			1	1	1		4
SR	B98-063	Strontium, Total in Water	SR8990	1			1	1	1		4
U	B98-063	Uranium, Isotopic in Water	UPLATE	1			1	1	1		4
TOTALS				5			5	5	5		20

### WORK SUMMARY

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### SUMMARY DATA SECTION

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Lab id TMANC

Protocol Hanford

Version Ver 1.0

Form DVD-CWS

Version 3.06

Report date 08/27/98

**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

**N808012-03**

**METHOD BLANK**

**Method Blank**

SDG <u>7486</u>	Client/Case no <u>Hanford</u>	SDG <u>H0186</u>
Contact <u>N. Joseph Verville</u>	Case no <u>TRB-SBB-207925</u>	
Lab sample id <u>N808012-03</u>	Client sample id <u>Method Blank</u>	
Dept sample id <u>7486-003</u>	Material/Matrix <u>LIQUID</u>	
	SAF No <u>B98-063</u>	

ANALYTE	CAS NO	RESULT pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST
Uranium 233/234	U-233/234	0	0.005	0.021	1.0	U	U
Uranium 235	15117-96-1	0	0.007	0.025	1.0	U	U
Uranium 238	U-238	0	0.005	0.021	1.0	U	U
Plutonium 238	13981-16-3	0.003	0.006	0.022	1.0	U	PU
Plutonium 239/240	15117-48-3	-0.003	0.006	0.022	1.0	U	PU
Americium 241	14596-10-2	0.007	0.011	0.019		U	AM
Total Strontium	SR-89/90	-0.022	0.15	0.21	2.0	U	SR
GAMMA SCAN ANALYTES		U					
Potassium 40	13966-00-2	U		230		U	GAM
Cobalt 60	10198-40-0	U		16	25	U	GAM
Cesium 137	10045-97-3	U		13	15	U	GAM
Europium 152	14683-23-9	U		35	50	U	GAM
Europium 154	15585-10-1	U		40	50	U	GAM
Europium 155	14391-16-3	U		34	50	U	GAM

QC-BLANK 28822

## TMA/RICHMOND

SAMPLE DELIVERY GROUP H0186

N808012-02

Lab Control Sample

## LAB CONTROL SAMPLE

SDG 7486  
Contact N. Joseph VervilleClient/Case no Hanford SDG H0186  
Case no TRB-SBB-207925Lab sample id N808012-02  
Dept sample id 7486-002Client sample id Lab Control Sample  
Material/Matrix LIQUID  
SAF No B98-063

ANALYTE	RESULT pCi/L	2 $\sigma$ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST	ADDED pCi/L	2 $\sigma$ ERR pCi/L	REC %	3 $\sigma$ LMITS (TOTAL)	PROTOCOL LIMITS
Uranium 233/234	5.1	0.33	0.15	1.0		U	5.14	0.21	99	86-114	80-120
Uranium 235	4.1	0.28	0.022	1.0		U	4.20	0.17	98	86-114	80-120
Uranium 238	5.4	0.34	0.14	1.0		U	5.29	0.21	102	86-114	80-120
Plutonium 238	5.7	0.53	0.023	1.0		PU	5.69	0.23	100	83-117	80-120
Plutonium 239/240	5.6	0.52	0.028	1.0		PU	5.29	0.21	106	82-118	80-120
Americium 241	5.2	0.30	0.016			AM	5.28	0.21	98	87-113	
Total Strontium	9.4	0.67	0.36	2.0		SR	9.89	0.40	95	81-119	
GAMMA SCAN ANALYTES	U										
Cobalt 60	650	49	23	25		GAM	584	23	111	71-129	80-120
Cesium 137	520	39	<u>30</u>	15		GAM	450	18	116	70-130	80-120

QC-LCS 28821

LAB CONTROL SAMPLES

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SUMMARY DATA SECTION

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Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-LCS  
Version 3.06  
Report date 08/27/98



## TMA/RICHMOND

SAMPLE DELIVERY GROUP H0186

N808012-04

B0NCD1

## DUPLICATE

SDG 7486

Contact N. Joseph Verville

DUPLICATE

Lab sample id N808012-04Dept sample id 7486-004

ORIGINAL

Lab sample id N808012-01Dept sample id 7486-001Received 08/05/98Client/Case no Hanford SDG H0186Case no TRB-SBB-207925Client sample id B0NCD1Location/Matrix 105-C LIQUIDCollected 08/03/98 16:10Custody/SAF No B98-063-05 B98-063

ANALYTE	DUPLICATE pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST	ORIGINAL pCi/L	2σ ERR (COUNT)	MDA pCi/L	QUALI- FIERS	RPD %	3σ PROT TOT LIMIT
Uranium 233/234	0.016	0.021	0.041	1.0	U	U	0.005	0.010	0.038	U	-	
Uranium 235	0.006	0.013	0.049	1.0	U	U	0	0.012	0.046	U	-	
Uranium 238	0	0.011	0.041	1.0	U	U	0	0.010	0.038	U	-	
Plutonium 238	-0.003	0.006	0.021	1.0	U	PU	-0.004	0.009	0.034	U	-	
Plutonium 239/240	0	0.006	0.021	1.0	U	PU	-0.004	0.009	0.034	U	-	
Americium 241	0.030	0.021	0.024			AM	0.016	0.012	0.015		61	158
Total Strontium	-0.064	0.36	0.46	2.0	U	SR	0.063	0.30	0.41	U	-	
GAMMA SCAN ANALYTES	U						U					
Potassium 40	U		78		U	GAM	U		78	U	-	
Cobalt 60	U		5.8	25	U	GAM	U		6.1	U	-	
Cesium 137	U		5.3	15	U	GAM	U		5.1	U	-	
Europium 152	U		17	50	U	GAM	U		15	U	-	
Europium 154	U		18	50	U	GAM	U		20	U	-	
Europium 155	U		12	50	U	GAM	U		12	U	-	

QC-DUP#1 28823

DUPLICATES

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SUMMARY DATA SECTION

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Lab id TMANCProtocol HanfordVersion Ver 1.0Form DVD-DUPVersion 3.06Report date 08/27/98

**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

**N808012-01**

**BONCD1**

**DATA SHEET**

SDG <u>7486</u>	Client/Case no <u>Hanford</u>	SDG <u>H0186</u>
Contact <u>N. Joseph Verville</u>	Case no <u>TRB-SBB-207925</u>	
Lab sample id <u>N808012-01</u>	Client sample id <u>BONCD1</u>	
Dept sample id <u>7486-001</u>	Location/Matrix <u>105-C</u>	<u>LIQUID</u>
Received <u>08/05/98</u>	Collected <u>08/03/98 16:10</u>	
	Custody/SAF No <u>B98-063-05</u>	<u>B98-063</u>

ANALYTE	CAS NO	RESULT pCi/L	2σ ERR (COUNT)	MDA pCi/L	RDL pCi/L	QUALI- FIERS	TEST
Uranium 233/234	U-233/234	0.005	0.010	0.038	1.0	U	U
Uranium 235	15117-96-1	0	0.012	0.046	1.0	U	U
Uranium 238	U-238	0	0.010	0.038	1.0	U	U
Plutonium 238	13981-16-3	-0.004	0.009	0.034	1.0	U	PU
Plutonium 239/240	15117-48-3	-0.004	0.009	0.034	1.0	U	PU
Americium 241	14596-10-2	0.016	0.012	0.015			AM
Total Strontium	SR-89/90	0.063	0.30	0.41	2.0	U	SR
GAMMA SCAN ANALYTES		U					
Potassium 40	13966-00-2	U		78		U	GAM
Cobalt 60	10198-40-0	U		6.1	25	U	GAM
Cesium 137	10045-97-3	U		5.1	15	U	GAM
Europium 152	14683-23-9	U		15	50	U	GAM
Europium 154	15585-10-1	U		20	50	U	GAM
Europium 155	14391-16-3	U		12	50	U	GAM

**TMA/RICHMOND**

SAMPLE DELIVERY GROUP H0186

Test AM Matrix LIQUIDSDG 7486Contact N. Joseph Verville**METHOD SUMMARY**

AMERICIUM 241 IN WATER

ALPHA SPECTROSCOPY

Client HanfordContract TRB-SBB-207925Case no SDG H0186**RESULTS**

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST FIX	SUF- PLANCHET	Americium 241
------------------	------------------	-----------------	------------------	------------------

Preparation batch 2785-103

BONCD1	N808012-01	7486-001	0.016
BLK (QC ID=28822)	N808012-03	7486-003	U
LCS (QC ID=28821)	N808012-02	7486-002	ok
Duplicate (N808012-01)	N808012-04	7486-004	ok

Nominal values and limits from method RDLs (pCi/L)

**METHOD PERFORMANCE**

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST FIX	SUF- pCi/L	MDA	ALIQ L	PREP FAC	DILU- TION	YIELD %	EFP %	COUNT min	FWHM keV	DRIFT KeV	DAYS HELD	ANAL- YZED	DETECTOR
------------------	------------------	-----------------	---------------	-----	-----------	-------------	---------------	------------	----------	--------------	-------------	--------------	--------------	---------------	----------

Preparation batch 2785-103 2σ prep error 5.0 % Reference Lab Notebook #2785 pg. 105

BONCD1	N808012-01	0.015	1.00	88	1087	22	08/21/98	08/25	SS-035
BLK (QC ID=28822)	N808012-03	0.019	1.00	78	1087	08/21/98	08/25	SS-038	
LCS (QC ID=28821)	N808012-02	0.016	1.00	83	1087	08/21/98	08/25	SS-036	
Duplicate (N808012-01)	N808012-04	0.024	1.00	42	1255	23	08/21/98	08/26	SS-043
(QC ID=28823)									

Nominal values and limits from method 1.00 20-105 700 100 180

PROCEDURES	REFERENCE	AM/CMPLATE
EP-040	Environmental Water Dissolution, rev 1	
EP-940	Plutonium Purification, rev 0	
EP-960	Americium-Curium Purification, rev 0	
EP-008	Heavy Elements Electroplating, rev 0	

AVERAGES ± 2 SD	MDA	0.018 ± 0.008
FOR 4 SAMPLES	YIELD	73 ± 42

**METHOD SUMMARIES**

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**SUMMARY DATA SECTION**

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Lab id	<u>TMANC</u>
Protocol	<u>Hanford</u>
Version	<u>Ver 1.0</u>
Form	<u>DVD-CMS</u>
Version	<u>3.06</u>
Report date	<u>08/27/98</u>

## TMA/RICHMOND

SAMPLE DELIVERY GROUP H0186

Test PU Matrix LIQUID

SDG 7486

Contact N. Joseph Verville

## METHOD SUMMARY

PLUTONIUM, ISOTOPIC IN WATER

ALPHA SPECTROSCOPY

Client Hanford

Contract TRB-SBB-207925

Case no SDG H0186

## RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST FIX	SUF- PLANCHET	Plutonium 238	Plutonium 239/240
------------------	------------------	-----------------	------------------	------------------	----------------------

## Preparation batch 2785-103

BONCD1	N808012-01	7486-001	U	U
BLK (QC ID=28822)	N808012-03	7486-003	U	U
LCS (QC ID=28821)	N808012-02	7486-002	ok	ok
Duplicate (N808012-01)	N808012-04	7486-004	- U	- U

Nominal values and limits from method	RDLs (pCi/L)	1.0	1.0
---------------------------------------	--------------	-----	-----

## METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST FIX	SUF- pCi/L	MAX MDA L	ALIQ L	PREP FAC	DILU- TION	YIELD %	EFF %	COUNT min	FWHM keV	DRIFT KeV	DAYS HELD	ANAL- YZED	DETECTOR
------------------	------------------	-----------------	---------------	--------------	-----------	-------------	---------------	------------	----------	--------------	-------------	--------------	--------------	---------------	----------

## Preparation batch 2785-103 2σ prep error 5.0 % Reference Lab Notebook #2785 pg. 105

BONCD1	N808012-01	0.034	1.00	61	619	16	08/19/98	08/19	SS-052
BLK (QC ID=28822)	N808012-03	0.022	1.00	65	619	08/19/98	08/19	SS-062	
LCS (QC ID=28821)	N808012-02	0.028	1.00	63	619	08/19/98	08/19	SS-058	
Duplicate (N808012-01)	N808012-04	0.021	1.00	67	619	16	08/19/98	08/19	SS-066
(QC ID=28823)									

Nominal values and limits from method	1.0	1.00	20-105	700	100	180
---------------------------------------	-----	------	--------	-----	-----	-----

PROCEDURES	REFERENCE	PUPLATE
EP-040	Environmental Water Dissolution, rev 1	
EP-940	Plutonium Purification, rev 0	
EP-008	Heavy Elements Electroplating, rev 0	

AVERAGES ± 2 SD	MDA	0.026 ± 0.012
FOR 4 SAMPLES	YIELD	64 ± 5

## METHOD SUMMARIES

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## SUMMARY DATA SECTION

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Lab id	TMANC
Protocol	Hanford
Version	Ver 1.0
Form	DVD-CMS
Version	3.06
Report date	08/27/98

# TMA/RICHMOND

SAMPLE DELIVERY GROUP H0186

Test U Matrix LIQUID  
SDG 7486  
Contact N. Joseph Verville

## METHOD SUMMARY

URANIUM, ISOTOPIC IN WATER  
ALPHA SPECTROSCOPY

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

## RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX	1: Uranium 233/234	2: Uranium 235	3: Uranium 238	RESULT RATIOS (%)			
			PLANCHET				1+3	2σ	2+3	2σ
Preparation batch 2785-103										
BONCD1	N808012-01		7486-001	U	U	U				
BLK (QC ID=28822)	N808012-03		7486-003	U	U	U				
LCS (QC ID=28821)	N808012-02		7486-002	ok	ok	ok				
Duplicate (N808012-01)	N808012-04		7486-004	- U	- U	- U				
Nominal values and limits from method										
			RDLs (pCi/L)	1.0	1.0	1.0	100			4
Averages										

## METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX	MAX MDA pCi/L	ALIQ L	PREP FAC	DILU- TION	YIELD %	EFF %	COUNT min	FWHM keV	DRIFT KeV	DAYS HELD	ANAL- PREPARED	YZED	DETECTOR
Preparation batch 2785-103 2σ prep error 5.0 % Reference Lab Notebook #2785 pg. 105																
BONCD1	N808012-01			0.046	0.500			89	525				10	08/13/98	08/13	SS-035
BLK (QC ID=28822)	N808012-03			0.025	1.00			84	525					08/13/98	08/13	SS-037
LCS (QC ID=28821)	N808012-02			0.15	1.00			94	525					08/13/98	08/13	SS-036
Duplicate (N808012-01)	N808012-04			0.049	0.500			86	525				10	08/13/98	08/13	SS-038
(QC ID=28823)																
Nominal values and limits from method																
				1.0	1.00			30-105	150	100			180			

PROCEDURES	REFERENCE	UPLATE
EP-040	Environmental Water Dissolution, rev 1	
EP-910	Uranium Purification, rev 0	
EP-008	Heavy Elements Electroplating, rev 0	

AVERAGES ± 2 SD	MDA <u>0.068</u> ± <u>0.11</u>
FOR 4 SAMPLES	YIELD <u>88</u> ± <u>9</u>

### METHOD SUMMARIES

Page 3

### SUMMARY DATA SECTION

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Lab id	<u>TMANC</u>
Protocol	<u>Hanford</u>
Version	<u>Ver 1.0</u>
Form	<u>DVD-CMS</u>
Version	<u>3.06</u>
Report date	<u>08/27/98</u>

## TMA/RICHMOND

SAMPLE DELIVERY GROUP H0186

Test SR Matrix LIQUIDSDG 7486Contact N. Joseph Verville

## METHOD SUMMARY

STRONTIUM, TOTAL IN WATER

BETA COUNTING

Client HanfordContract TRB-SBB-207925Case no SDG H0186

## RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX	Total Strontium
------------------	------------------	-------------	-------------	--------------------

Preparation batch 2785-103

BONCD1	N808012-01	7486-001	U
BLK (QC ID=28822)	N808012-03	7486-003	U
LCS (QC ID=28821)	N808012-02	7486-002	ok
Duplicate (N808012-01)	N808012-04	7486-004	- U

Nominal values and limits from method RDLs (pCi/L) 2.0

## METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST	SUF- FIX	MDA pCi/L	ALIQ L	PREP FAC	DILU- TION	YIELD %	EFF %	COUNT min	FWHM keV	DRIFT KeV	DAYS HELD	ANAL- YZED	DETECTOR
------------------	------------------	-------------	-------------	--------------	-----------	-------------	---------------	------------	----------	--------------	-------------	--------------	--------------	---------------	----------

Preparation batch 2785-103 2σ prep error 10.0 % Reference Lab Notebook #2785 pg. 105

BONCD1	N808012-01	0.41	<u>0.500</u>	79	400	10	08/13/98	08/13	GRB-205
BLK (QC ID=28822)	N808012-03	0.21	1.00	80	400	08/13/98	08/13	GRB-207	
LCS (QC ID=28821)	N808012-02	0.36	1.00	78	400	08/13/98	08/13	GRB-206	
Duplicate (N808012-01)	N808012-04	0.46	<u>0.500</u>	78	400	10	08/13/98	08/13	GRB-208
(QC ID=28823)									

Nominal values and limits from method 2.0 1.00 100 180

PROCEDURES	REFERENCE	SR8990
EP-040	Environmental Water Dissolution, rev 1	
EP-500	Strontium-89,90 - Purification, rev 0	
EP-519	Strontium-89,90 Planchet Demounting and Yttrium Purification, rev 0	

AVERAGES ± 2 SD	MDA	<u>0.36</u> ± <u>0.22</u>
FOR 4 SAMPLES	YIELD	<u>79</u> ± <u>2</u>

## METHOD SUMMARIES

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## SUMMARY DATA SECTION

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Lab id	<u>TMANC</u>
Protocol	<u>Hanford</u>
Version	<u>Ver 1.0</u>
Form	<u>DVD-CMS</u>
Version	<u>3.06</u>
Report date	<u>08/27/98</u>

## TMA/RICHMOND

SAMPLE DELIVERY GROUP H0186

Test GAM Matrix LIQUIDSDG 7486Contact N. Joseph Verville

## METHOD SUMMARY

GAMMA EMITTERS

GAMMA SCAN

Client HanfordContract TRB-SBB-207925Case no SDG H0186

## RESULTS

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST FIX	SUF- PLANCHET	Cobalt 60	Cesium 137
------------------	------------------	-----------------	------------------	-----------	------------

Preparation batch 2785-103

BONCD1	N808012-01	7486-001	U	U	
BLK (QC ID=28822)	N808012-03	7486-003	U	U	
LCS (QC ID=28821)	N808012-02	7486-002	ok	ok	
Duplicate (N808012-01)	N808012-04	7486-004	-	U	- U

Nominal values and limits from method	RDLs (pCi/L)	25	15
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## METHOD PERFORMANCE

CLIENT SAMPLE ID	LAB SAMPLE ID	RAW TEST FIX	SUF- pCi/L	MAX MDA L	ALIQ L	PREP FAC	DILU- TION	YIELD %	EFF %	COUNT min	FWHM keV	DRIFT KeV	DAYS HELD	ANAL- PREPARED	YZED	DETECTOR
------------------	------------------	-----------------	---------------	--------------	-----------	-------------	---------------	------------	----------	--------------	-------------	--------------	--------------	-------------------	------	----------

Preparation batch 2785-103 2σ prep error 15.0 % Reference Lab Notebook #2785 pg. 105

BONCD1	N808012-01	5.1	0.500	516	8	08/11/98	08/11	01,04,00								
BLK (QC ID=28822)	N808012-03	13	0.500	455	8	08/11/98	08/11	01,03,00								
LCS (QC ID=28821)	N808012-02	30	0.500	455	8	08/11/98	08/11	01,01,00								
Duplicate (N808012-01)	N808012-04	5.3	0.500	505	9	08/11/98	08/12	01,04,00								
(QC ID=28823)																

Nominal values and limits from method	15	0.500	400	180
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PROCEDURES REFERENCE GAMMAHI

EP-100 Ge(Li) Preparation for Environmental Samples,  
rev 0

AVERAGES ± 2 SD

FOR 4 SAMPLES

MDA 13 ± 23YIELD        ±       

METHOD SUMMARIES

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SUMMARY DATA SECTION

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Lab id TMANCProtocol HanfordVersion Ver 1.0Form DVD-QMSVersion 3.06Report date 08/27/98

**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**REPORT GUIDE**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**DATA SHEET**

The Data Sheet Report shows all results and primary supporting information for one client sample or Method Blank. This report corresponds to both the CLP Inorganics and Organics Data Sheet.

The following notes apply to this report:

- \* TEST is a code for the method used to measure an analyte. If the TEST is empty, no data is available; the analyte was not analyzed for.
- \* The LAB SAMPLE ID and TEST uniquely identify work within the Summary Data Section of a Data Package. The Work Summary and Method Summary Reports further identify raw data that underlies this work.

The Method Summary Report for each TEST has method performance data, such as yield, for each Lab Sample ID and a list of procedures used in the method.

- \* ERRORS can be labeled TOTAL or COUNT. TOTAL implies a preparation (non-counting method) error has been added, as square root of sum of squares, to the counting error denoted by COUNT. The preparation errors, which may vary by preparation batch, are shown on the Method Summary Report.
- \* A RESULT can be 'N.R.' (Not Reported). This means the lab did this work but chooses not to report it now, possibly because it was reported at another time.
- \* When reporting a Method Blank, a RESULT can be 'N.A.' (Not Applicable). This means there is no reported client sample work in the same preparation batch as the Blank's result. This is likely to occur when the Method Blank is associated with reanalyses of selected work for a few samples in the SDG.

The following qualifiers are defined by the DVD system:

- U The RESULT is less than the MDA (Minimum Detectable Activity).

**REPORT GUIDES**

Page 4

**SUMMARY DATA SECTION**

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Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98



**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**GUIDE, cont.**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**DATA SHEET**

If the MDA is blank, the ERROR is used as the limit.

- J The RESULT is less than the RDL (Required Detection Limit) and no U qualifier is assigned.
- B A Method Blank associated with this sample had a result without a U flag and, after correcting for possibly different aliquots, that result is greater than or equal to the MDA for this sample.

Normally, B is not assigned if U is. When method blank subtraction is shown on this report, B flags are assigned based on the unsubtracted values while U's are assigned based on the subtracted ones. Both flags can be assigned in this case.

For each sample result, all Method Blank results in the same preparation batch are compared. The Method Summary Report documents this and other QC relationships.

- L Some Lab Control Sample that QC's this sample had a low recovery. The lab can disable assignment of this qualifier.
- H Similar to 'L' except the recovery was high.
- P The RESULT is 'preliminary'.
- X Some data necessary to compute the RESULT, ERROR or MDA was manually entered or modified.
- 2 There were two or more results available for this analyte. The reported result may not be the same as in the raw data.

Other qualifiers are lab defined. Definitions should be in the SDG narrative.

The following values are underlined to indicate possible problems:

- \* An MDA is underlined if it is bigger than its RDL.

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

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Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98

**TMA/RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**GUIDE, cont.**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**DATA SHEET**

- \* An ERROR is underlined if the 1.645 sigma counting error is bigger than both the MDA and the RESULT, implying that the MDA may not be a good estimate of the 'real' minimum detectable activity.
- \* A negative RESULT is underlined if it is less than the negative of its 2 sigma counting ERROR.
- \* When reporting a Method Blank, a RESULT is underlined if greater than its MDA. If the MDA is blank, the 2 sigma counting error is used in the comparison.

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

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Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98

**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**REPORT GUIDE**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**LAB CONTROL SAMPLE**

The Lab Control Sample Report shows all results, recoveries and primary supporting information for one Lab Control Sample.

The following notes apply to this report:

- \* All fields in common with the Data Sheet Report have similar usage. Refer to its Report Guide for details.
- \* An amount ADDED is the lab's value for the actual amount spiked into this sample with its ERROR an estimate of the error of this amount.

An amount added is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- \* REC (Recovery) is RESULT divided by ADDED expressed as a percent.
- \* The first, computed limits for the recovery reflect:
  1. The error of RESULT, including that introduced by rounding the result prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.
  2. The error of ADDED.
  3. A lab specified, per analyte bias. The bias changes the center of the computed limits.
- \* The second limits are protocol defined upper and lower QC limits for the recovery.
- \* The recovery is underlined if it is outside either of these ranges.

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

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Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
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**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**REPORT GUIDE**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**DUPLICATE**

The Duplicate Report shows all results, differences and primary supporting information for one Duplicate and associated Original sample.

The following notes apply to this report:

- \* All fields in common with the Data Sheet Report have similar usage. This applies both to the Duplicate and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Duplicate has data for a TEST and the lab did not do this test to the Original, the Original's RESULTS are underlined.

- \* The RPD (Relative Percent Difference) is the absolute value of the difference of the RESULTS divided by their average expressed as a percent.

If both RESULTS are less than their MDAs, no RPD is computed and a '-' is printed.

For an analyte, if the lab did work for both samples but has data for only one, the MDA from the sample with data is used as the other's result in the RPD.

- \* The first, computed limit is the sum, as square root of sum of squares, of the errors of the results divided by the average result as a percent, hence the relative error of the difference rather than the error of the relative difference. The errors include those introduced by rounding the RESULTS prior to printing.

If this limit is labeled TOT, it includes the preparation error in the RESULTS. If labeled CNT, it does not.

This value reported for this limit is at most 999.

- \* The second limit for the RPD is the larger of:
  1. A fixed percentage specified in the protocol.

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

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Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
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**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**GUIDE, cont.**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**DUPLICATE**

2. A protocol factor (typically 2) times the average MDA as a percent of the average result. This limit applies when the results are close to the MDAs.

- \* The RPD is underlined if it is greater than either limit.
- \* If specified by the lab, the second limit column is replaced by the Difference Error Ratio (DER), which is the absolute value of the difference of the results divided by the quadratic sum of their one sigma errors, the same errors as used in the first limit.

Except for differences due to rounding, the DER is the same as the RPD divided by the first RPD limit with the limit scaled to 1 sigma.

- \* The DER is underlined if it is greater than the sigma factor, typically 2 or 3, shown in the header for the first RPD limit.

Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98

**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**REPORT GUIDE**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**MATRIX SPIKE**

The Matrix Spike Report shows all results, recoveries and primary supporting information for one Matrix Spike and associated Original sample.

The following notes apply to this report:

- \* All fields in common with the Data Sheet Report have similar usage. This applies both to the Spiked and Original sample data. Refer to the Data Sheet Report Guide for details.

If the Spike has data for a TEST and the lab did not do this test to the Original, the Original's RESULTS are underlined.

- \* An amount ADDED is the lab's value for the actual amount spiked into the Spike sample with its ERROR an estimate of the error of this amount.

An amount is underlined if its ratio to the corresponding RDL is outside protocol specified limits.

- \* REC (Recovery) is the Spike RESULT minus the Original RESULT divided by ADDED expressed as a percent.

- \* The first, computed limits for the recovery reflect:

1. The errors of the two RESULTS, including those introduced by rounding them prior to printing.

If the limits are labeled (TOTAL), they include preparation error in the result. If labeled (COUNT), they do not.

2. The error of ADDED.

3. A lab specified, per analyte bias. The bias changes the center of the computed limits.

- \* The second limits are protocol defined upper and lower QC limits

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

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Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98

**TMA/RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**GUIDE, cont.**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**MATRIX SPIKE**

for the recovery.

These limits are left blank if the Original RESULT is more than a protocol defined factor (typically 4) times ADDED. This is a way of accounting for that when the spike is small compared to the amount in the original sample, the recovery is unreliable.

- \* The recovery is underlined (out of spec) if it is outside either of these ranges.

**REPORT GUIDES**

Page 11

**SUMMARY DATA SECTION**

Page 26

Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98

**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**REPORT GUIDE**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**METHOD SUMMARY**

The Method Summary Report has two tables. One shows up to five results measured using one method. The other has performance data for the method. There is one report for each TEST, as used on the Data Sheet Report.

The following notes apply to this report:

- \* Each table is subdivided into sections, one for each preparation batch. A preparation batch is a group of aliquots prepared at roughly the same time in one work area of the lab using the same method.

There should be Lab Control Sample and Method Blank results in each preparation batch since this close correspondence makes the QC meaningful. Depending on lab policy, Duplicates need not occur in each batch since they QC sample dependencies such as matrix effects.

- \* The RAW TEST column shows the test code used in the raw data to identify a particular analysis if it is different than the test code in the header of the report. This occurs in special cases due to method specific details about how the lab labels work.

The Lab Sample or Planchet ID combined with the (Raw) Test Code and Suffix uniquely identify the raw data for each analysis.

- \* If a result is less than both its MDA and RDL, it is replaced by just 'U' on this report. If it is greater than or equal to the RDL but less than the MDA, the result is shown with a 'U' flag.

The J and X flags are as on the data sheet.

- \* Non-U results for Method Blanks are underlined to indicate possible contamination of other samples in the preparation batch. The Method Blank Report has supporting data.
- \* Lab Control Sample and Matrix Spike results are shown as: ok, No data, LOW or HIGH, with the last two underlined. 'No data'

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

Page 27

Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98



**TMA/RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**GUIDE, cont.**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**METHOD SUMMARY**

means no amount ADDED was specified. 'LOW' and 'HIGH' correspond to when the recovery is underlined on the Lab Control Sample or Matrix Spike Report. See these reports for supporting data.

- \* Duplicate sample results are shown as: ok, No data, or OUT, with the last two underlined. 'No data' means there was no original sample data found for this duplicate. 'OUT' corresponds to when the RPD is underlined on the Duplicate Report. See this report for supporting data.
- \* If the MDA column is labeled 'MAX MDA', there was more than one result measured by the reported method and the MDA shown is the largest MDA. If not all these results have the same RDL, the MAX MDA reflects only those results with RDL equal to the smallest one.

MDAs are underlined if greater than the printed RDL.

- \* Aliquots are underlined if less than the nominal value specified for the method.
- \* Preparation factors are underlined if greater than the nominal value specified for the method.
- \* Dilution factors are underlined if greater than the nominal value specified for the method.
- \* Residues are underlined if outside the range specified for the method. Residues are not printed if yields are.
- \* Yields, which may be gravimetric, radiometric or some type of recovery depending on the method, are underlined if outside the range specified for the method.
- \* Efficiencies are underlined if outside the range specified for the method. Efficiencies are detector and geometry dependent so this test is only approximate.

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

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Lab id TMANC  
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**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**GUIDE, cont.**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**METHOD SUMMARY**

- \* Count times are underlined if less than the nominal value specified for the method.
- \* Resolutions (as FWHM; Full Width at Half Max) are underlined if greater than the method specified limit.
- \* Tracer drifts are underlined if their absolute values are greater than the method specified limit. Tracer drifts are not printed if percent moistures are.
- \* Days Held are underlined if greater than the holding time specified in the protocol.
- \* Analysis dates are underlined if before their planchet's preparation date or, if a limit is specified, too far after it.

For some methods, ratios as percentages and error estimates for them are computed for pairs of results. A ratio column header like '1+3' means the ratio of the first result column and the third result column.

Ratios are not computed for Lab Control Sample, Method Blank or Matrix Spike results since their matrices are not necessarily similar to client samples'.

The error estimate for a ratio of results from one planchet reflects only counting errors since other errors should be correlated. For a ratio involving different planchets, if QC limits are computed based on total errors, the error for the ratio allows for the preparation errors for the planchets.

The ratio is underlined (out of spec) if the absolute value of its difference from the nominal value is greater than its error estimate. If no nominal value is specified, this test is not done.

For Gross Alpha or Gross Beta results, there may be a column showing the sum of other Alpha or Beta emitters. This sum includes all relevant

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

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Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98

**TMA / RICHMOND**  
**SAMPLE DELIVERY GROUP H0186**

SDG 7486  
Contact N. Joseph Verville

**GUIDE, cont.**

Client Hanford  
Contract TRB-SBB-207925  
Case no SDG H0186

**METHOD SUMMARY**

results in the DVD database, whether reported or not. Results in the sum are weighted by a particles/decay value specified by the lab for each relevant analyte. Results less than their MDA are not included. No sums are computed for Lab Control, Method Blank or Matrix Spike samples since their various planchets may not be physically related.

If a ratio of total isotopic to Gross Alpha or Beta is shown, the error for the ratio reflects both the error in the Gross result and the sum, as square root of sum of squares, of the errors in the isotopic results.

For total elemental uranium or thorium results, there may be a column showing the total weight computed from associated isotopic results. Ignoring results less than their MDAs, this is a weighted sum of the isotopic results. The weights depend on the molecular weight and half-life of each isotope so as to convert activities (decays) to weight (atoms).

If a ratio of total computed to measured elemental uranium or thorium is shown, the error for the ratio reflects the errors in all the measurements.

**REPORT GUIDES**

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**SUMMARY DATA SECTION**

Page 30

Lab id TMANC  
Protocol Hanford  
Version Ver 1.0  
Form DVD-RG  
Version 3.06  
Report date 08/27/98

<b>Bechtel Hanford Inc.</b>		<b>CHAIN OF CUSTODY/SAMPLE ANALYSIS REQUEST</b>						<b>B98-063-05</b>		Page <b>1</b> of <b>2</b>	
Collector <b>Fahlberg/Nielson</b>		Company Contact <b>Steve Marske</b>		Telephone No. <b>373-4316</b>		Project Coordinator <b>WEISS, RL</b>		<b>Data Turnaround</b>  <b>21 Days</b>			
Project Designation <b>105-C Phase I &amp; Phase II - Water Samples</b>		Sampling Location <b>105-C</b>		SAF No. <b>B98-063</b>							
Ice Chest No.		Field Logbook No. <b>EL-1309-1</b>		Method of Shipment <b>Federal Express</b>							
Shipped To <b>RUN Thermo NUTech</b> <del>Quanterra Incorporated</del> <b>8/4/98 Richmond Laboratory</b>		Offsite Property No.		Bill of Lading/Air Bill No.							
Waste Designation				COA							

POSSIBLE SAMPLE HAZARDS/REMARKS	Preservation	None	HNO3 to pH <2	Cool 4C	HNO3 to pH <2	HNO3 to pH <2	HNO3 to pH <2	Cool 4C	HNO3 to pH <2	HNO3 to pH <2	HNO3 to pH <2
	Type of Container	P	aG	P	P	P	P	aG	P	P	P
	No. of Container(s)	1 ✓	1	1 ✓	1	1	1	2	3 <del>2</del> <b>3</b>	2 ✓	3
	Special Handling and/or Storage	Cool 4C									

SAMPLE ANALYSIS				Activity Scan	Mercury - 7470 - (CV)	Chromium Hex - 7196	ICP Metals - 6010A (Supertrace) (Lead)	See item (1) in Special Instructions	Nickel-63	PCBs - 8080 (Aroclor-1254)	See item (2) in Special Instructions	Isotopic Plutonium, Isotopic Uranium, Americium-241	Selenium-79
-----------------	--	--	--	---------------	-----------------------	---------------------	--	--------------------------------------	-----------	----------------------------	--------------------------------------	---	-------------

Sample No.	Matrix *	Sample Date	Sample Time										
B0NCD1	Water	8-3-98	16:0	X					X		X	X	X

CHAIN OF POSSESSION	Sign/Print Names		SPECIAL INSTRUCTIONS		Matrix *
Relinquished By <b>R. Nielson</b>	Date/Time <b>8/4/98</b>	Received By <b>Fed Ex</b>	Date/Time	<b>(1) Gamma Spectroscopy (Cesium-137, Cobalt-60, Europium-152, Europium-154, Europium-155)</b> <b>(2) Gamma Spectroscopy (Water) (Cesium-137, Cobalt-60, Europium-152, Europium-154, Europium-155)</b> <b>DO NOT run Nickel-63, Carbon-14, or Technetium-99 unless instructed by ERC Sample Management. ALSO DO NOT run Selenium-79 unless directed by ERC Sample Management.</b>	S = Soil SE = Sediment SO = Solid SL = Sludge W = Water O = Oil A = Air DS = Drum Solids DL = Drum Liquids T = Tissue WI = Wipe L = Liquid V = Vegetation X = Other
Relinquished By	Date/Time	Received By <b>R. SANGALANG</b>	Date/Time <b>8/5/98 1000</b>		
Relinquished By	Date/Time	Received By	Date/Time		
Relinquished By	Date/Time	Received By	Date/Time		

LABORATORY SECTION	Received By	Title	Date/Time
FINAL SAMPLE DISPOSITION	Disposal Method	Disposed By	Date/Time

<b>Bechtel Hanford Inc.</b>		<b>CHAIN OF CUSTODY/SAMPLE ANALYSIS REQUEST</b>					<b>B98-063-05</b>		Page <u>2</u> of <u>2</u>					
Collector Fahlberg/Nielson		Company Contact Steve Marske		Telephone No. 373-4316		Project Coordinator WEISS, RL		<b>Data Turnaround</b>  <b>21 Days</b>						
Project Designation 105-C Phase I & Phase II - Water Samples		Sampling Location 105-C		SAF No. B98-063										
Ice Chest No.		Field Logbook No. EL-1309-1		Method of Shipment Federal Express										
Shipped To <u>R/N Thermo Nutech</u> <u>Quanterra Incorporated</u> <u>8/4/98 Richmond Laboratory</u>		Offsite Property No.		Bill of Lading/Air Bill No.										
Waste Designation				COA										
<b>POSSIBLE SAMPLE HAZARDS/REMARKS</b>     <b>Special Handling and/or Storage</b> Cool 4C				<b>Preservation</b>		None	HCl to pH <2	HNO3 to pH <2						
				<b>Type of Container</b>		P	P	P						
				<b>No. of Container(s)</b>		3 ✓	3 ✓	3 ✓						
				<b>Volume</b>		500ml	1000ml	1000ml						
<b>SAMPLE ANALYSIS</b>				Carbon-14	Technetium-99	Strontium-89,90 -- Total Sr								
Sample No.	Matrix *	Sample Date	Sample Time											
BONCD1	Water	8-3-98	1610	X	X	X							BOP & Y2	
<b>CHAIN OF POSSESSION</b>		<b>Sign/Print Names</b>				<b>SPECIAL INSTRUCTIONS</b>  Do not run Nickel-63, Carbon-14, or Technetium-99 unless instructed by ERC Sample Management. ALSO do not run Selenium-79 unless directed by ERC Sample Management.					<b>Matrix *</b> S = Soil SE = Sediment SO = Solid SL = Sludge W = Water O = Oil A = Air DS = Drum Solids DL = Drum Liquids T = Tissue WI = Wipe L = Liquid V = Vegetation X = Other			
Relinquished By <u>Reene Nielson</u> <u>P Nielson</u> <u>8/4/98</u>		Received By <u>Fed Ex</u>												
Relinquished By		Received By <u>C. SANGALANG</u> <u>8/5/98</u>		<u>1000</u>										
Relinquished By		Received By												
Relinquished By		Received By												
<b>LABORATORY SECTION</b>		Received By				Title				Date/Time				
<b>FINAL SAMPLE DISPOSITION</b>		Disposal Method				Disposed By				Date/Time				

Figure 1

## SAMPLE CHECK-IN LIST

Date/Time Received: 8/5/98 / 1000 S.G.#: \_\_\_\_\_  
 Work Order Number: N8-08-11-12 SAF #: \_\_\_\_\_  
 Shipping Container ID: ERC 044 Chain of Custody #: B98-063-05 P-1 22

1. Custody Seals on shipping container intact? Yes [☒] No [☐]
2. Custody Seals dated and signed? Yes [☒] No [☐]
3. Chain-of-Custody record present? Yes [☒] No [☐]
4. Cooler temperature \_\_\_\_\_
5. Vermiculite/packing materials is Wet [☐] Dry [☐]
6. Number of samples in shipping container: 1 (20 bottles (poly))
7. Sample holding times exceeded? Yes [☐] No [☐]

8. Samples have:  
☒ tape ☐ hazard labels  
☒ custody seals ☒ appropriate sample labels

9. Samples are:  
☒ in good condition ☐ leaking  
☐ broken ☐ have air bubbles

10. Where any anomalies identified in sample receipt? Yes [☐] No [☒]

11. Description of anomalies (include sample numbers): There is a  
bottle for Chromium Hex-7196, but not  
requested for analysis in the chain of custody.  
Ice chest is not packed w/ ice.

Sample Custodian/Laboratory: university/TNH Date: 8/5/98

Telephoned To: \_\_\_\_\_ On \_\_\_\_\_ By \_\_\_\_\_

# Sample Disposition Record

Control #: B98-070

Revision#: 0

Date Initiated: 8/6/98

## Section 1 - BACKGROUND

SAF#: B98-063

OU: N/A

Project ID: 105-C Reactor

Task ID: 9

Sampling Event: 105-C Phase I & Phase II

Laboratory: TMA/RECRA

Project Coordinator: WEISS, RL

Task Manager: ARMATROUT, JF

## Section 2 - SAMPLE INFORMATION

Number of Samples: 1

ID Numbers: B0NCD1

MATRIX: Water

Collection Date:

## Section 3 - ISSUE

Class: Lab Direction

NCR Number: N/A

Type: Revision of Direction - Cancellation of Analyses

Description: Inadvertent inclusion of additional analytes to COC

N/A

NCR Validation (Print/Sign)

Date

## Section 4 - DISPOSITION

Type: Use As Is

Description: Cancel analysis for CrVI, Ni-63, C-14, and Tc-99

WEISS, RL



8/6/98

Project Coordinator (Print/Sign)

Date

ARMATROUT, JF

Task Manager (Print/Sign)

Date

N/A

QA (Print/Sign)

Date

## Section 5 - INSPECTION (Issue Class: Nonconformance Only)

Inspection Number: N/A

Inspection Results: N/A

N/A

Inspector (Print/Sign)

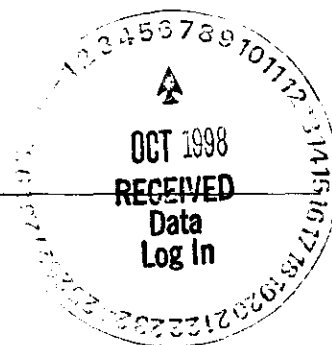
Date



**RECRA  
LabNet**

*a division of Recra Environmental, Inc.*

*Virtual Laboratories Everywhere*



**Recra LabNet Philadelphia  
Analytical Report**

**Client :** TNU-HANFORD B98-063

**RFW# :** 9808L205

**SDG/SAF# :** H0186/B98-063

**W.O.# :** 10985-001-001-9999-00

**Date Received:** 08-05-98

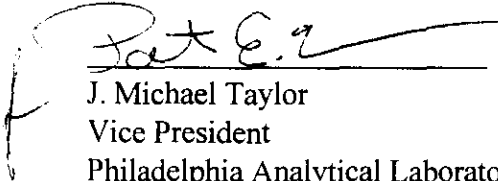
**METALS CASE NARRATIVE**

1. This narrative covers the analyses of 1 water sample.
2. The sample was prepared and analyzed in accordance with methods checked on the attached glossary.
3. All analyses were performed within the required holding times.
4. The cooler temperature has been recorded on the Chain of Custody.
5. All Initial and Continuing Calibration Verifications (ICV/CCVs) were within control limits.
6. All Initial and Continuing Calibration Blanks (ICB/CCBs) were within control limits.
7. All preparation/method blanks were within method criteria. Refer to the Inorganics Method Blank Data Summary.
8. All ICP Interference Check Standards were within control limits.
9. All laboratory control samples (LCS) were within the laboratory control limits. Refer to the Inorganics Laboratory Control Standards Report.
10. The matrix spike (MS) and matrix spike duplicate (MSD) recoveries for Lead were within the 75-125% control limits. Matrix QC could not be reported for Mercury. QC had been prepped, however, the rack of tubes with the replicate and spike was dropped and the tubes containing these samples broke. At this point, sample volume had been depleted. Refer to the Inorganics Accuracy Report.
11. The MSs and MSDs for Lead were within the 20% Relative Percent Difference (RPD) control limits. Refer to the Inorganics Matrix Spike Duplicate Report.
12. The duplicate analyses for Lead was outside the 20% Relative Percent Difference (RPD) control limits. Refer to the Inorganics Precision Report.

The results presented in this report relate only to the analytical testing and conditions of the samples at receipt and during storage. All pages of this report are integral parts of the analytical data. Therefore, this report should only be reproduced in its entirety of 13 pages.



13. For the purposes of this report, the data has been reported to the Instrument Detection Limit (IDL). Values between the IDL and the Practical Quantitation Limit (PQL) are acquired in a region of less-certain quantification.

  
J. Michael Taylor  
Vice President  
Philadelphia Analytical Laboratory

mld/m08-205

9-29-98  
Date



# METALS METHOD GLOSSARY

The following methods are used as reference for the digestion and analysis of samples contained within this Recra Lot#: 98084705

Leaching Procedure: 1310 1311 1312 Other: \_\_\_\_\_

CLP Metals Digestion and Analysis Methods: ILM03.0 ILM04.0

Metals Digestion Methods: X 3005A 3010A 3015 3020A 3050A 3051 200.7 SS17  
Other: \_\_\_\_\_

## Metals Analysis Methods

	SW846	EPA	STD MTD	EPA OSWR	USATHAMA
Aluminum	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Antimony	<u>6010B</u> <u>7041</u> <sup>5</sup>	<u>200.7</u> <u>204.2</u>			<u>99</u>
Arsenic	<u>6010B</u> <u>7060A</u> <sup>5</sup>	<u>200.7</u> <u>206.2</u>	<u>3113B</u>		<u>99</u>
Barium	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Beryllium	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Bismuth	<u>6010B</u> <sup>1</sup>	<u>200.7</u> <sup>1</sup>		<u>1620</u>	<u>99</u>
Boron	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Cadmium	<u>6010B</u> <u>7131A</u> <sup>5</sup>	<u>200.7</u> <u>213.2</u>			<u>99</u>
Calcium	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Chromium	<u>6010B</u> <u>7191</u> <sup>5</sup>	<u>200.7</u> <u>218.2</u>			<u>SS17</u>
Cobalt	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Copper	<u>6010B</u> <u>7211</u> <sup>5</sup>	<u>200.7</u> <u>220.2</u>			<u>99</u>
Iron	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Lead	<u>X</u> <u>6010B</u> <u>7421</u> <sup>5</sup>	<u>200.7</u> <u>239.2</u>	<u>3113B</u>		<u>99</u>
Lithium	<u>6010B</u> <u>7430</u> <sup>4</sup>	<u>200.7</u>		<u>1620</u>	<u>99</u>
Magnesium	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Manganese	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Mercury	<u>X</u> <u>7470A</u> <sup>3</sup> <u>7471A</u> <sup>3</sup>	<u>245.1</u> <sup>2</sup> <u>245.5</u> <sup>2</sup>			<u>99</u>
Molybdenum	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Nickel	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Potassium	<u>6010B</u> <u>7610</u> <sup>4</sup>	<u>200.7</u> <u>258.1</u> <sup>4</sup>			<u>99</u>
Rare Earths	<u>6010B</u> <sup>1</sup>	<u>200.7</u> <sup>1</sup>		<u>1620</u>	<u>99</u>
Selenium	<u>6010B</u> <u>7740</u> <sup>5</sup>	<u>200.7</u> <u>270.2</u>	<u>3113B</u>		<u>99</u>
Silicon	<u>6010B</u> <sup>1</sup>	<u>200.7</u>		<u>1620</u>	<u>99</u>
Silica	<u>6010B</u>	<u>200.7</u>		<u>1620</u>	<u>99</u>
Silver	<u>6010B</u> <u>7761</u> <sup>5</sup>	<u>200.7</u> <u>272.2</u>			<u>99</u>
Sodium	<u>6010B</u> <u>7770</u> <sup>4</sup>	<u>200.7</u> <u>273.1</u> <sup>4</sup>			<u>99</u>
Strontium	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Thallium	<u>6010B</u> <u>7841</u> <sup>5</sup>	<u>200.7</u> <u>279.2</u> <u>200.9</u>			<u>99</u>
Tin	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Titanium	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Uranium	<u>6010B</u> <sup>1</sup>	<u>200.7</u> <sup>1</sup>		<u>1620</u>	<u>99</u>
Vanadium	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Zinc	<u>6010B</u>	<u>200.7</u>			<u>99</u>
Zirconium	<u>6010B</u> <sup>1</sup>	<u>200.7</u> <sup>1</sup>		<u>1620</u>	<u>99</u>

Other: \_\_\_\_\_

Method: \_\_\_\_\_

## **METHOD REFERENCES AND DATA QUALIFIERS**

### **DATA QUALIFIERS**

- U = Indicates that the parameter was not detected at or above the reported limit. The associated numerical value is the sample detection limit.
- \* = Indicates that the original sample result is greater than 4x the spike amount added.

### **ABBREVIATIONS**

- MB = Method or Preparation Blank.  
MS = Matrix Spike.  
MSD = Matrix Spike Duplicate.  
REP = Sample Replicate  
LCS = Laboratory Control Sample.  
NC = Not calculated.

### **ANALYTICAL METAL METHODS**

1. Not included in the method element list.
2. Modified Hg: Hg1 and Hg2 require less total volume of digestate due to the autosampler analysis. Sample volumes and reagents for mercury determinations in water and soil have been proportionately scaled down to adapt to this semi-automated technique. The sample volume used for water analysis is 33 mL. For soils, 0.1 grams of sample is taken to a final volume of 50 mL (including all reagents).
3. Modified Hg: Hg1 and Hg2 require less total volume of digestate due to the autosampler analysis. Sample volumes and reagents for mercury determinations in water and soil have been proportionately scaled down to adapt to this semi-automated technique. The sample volume used for water analysis is 33 mL. For soils, three 0.1 gram of sample is taken to a final volume of 50 mL (including all reagents).
4. Flame AA.
5. Graphite Furnace AA.

Recra LabNet - Lionville

INORGANICS DATA SUMMARY REPORT 09/29/98

CLIENT: TNU-HANFORD B98-063

RECRA LOT #: 9808L205

WORK ORDER: 10985-001-001-9999-00

SAMPLE	SITE ID	ANALYTE	RESULT	UNITS	REPORTING LIMIT	DILUTION FACTOR
-001	BONCD1	Mercury, Total	0.10 u	UG/L	0.10	1.0
		Lead, Total	4.8	UG/L	1.9	1.0

Recra LabNet - Lionville

INORGANICS METHOD BLANK DATA SUMMARY PAGE 09/29/98

CLIENT: TNU-HANFORD B98-063

RECRA LOT #: 9808L205

WORK ORDER: 10985-001-001-9999-00

SAMPLE	SITE ID	ANALYTE	RESULT	UNITS	REPORTING LIMIT	DILUTION FACTOR
BLANK1	98C0408-MB1	Mercury, Total	0.10 u	UG/L	0.10	1.0
BLANK1	98L1125-MB1	Lead, Total	1.9 u	UG/L	1.9	1.0

Recra LabNet - Lionville

INORGANICS ACCURACY REPORT 09/29/98

CLIENT: TNU-HANFORD B98-063

RECRA LOT #: 9808L205

WORK ORDER: 10985-001-001-9999-00

SAMPLE	SITE ID	ANALYTE	SPIKED SAMPLE	INITIAL RESULT	SPIKED AMOUNT	%RECOV	DILUTION FACTOR (SPK)
-001	BONCD1	Lead, Total	502	4.8	500	99.3	1.0
		Lead, Total MSD	489	4.8	500	96.9	1.0

Recra LabNet - Lionville

INORGANICS DUPLICATE SPIKE REPORT 09/29/98

CLIENT: TNU-HANFORD B98-063

RECRA LOT #: 9808L205

WORK ORDER: 10985-001-001-9999-00

SAMPLE	SITE ID	ANALYTE	SPIKE#1 SPIKE#2		
			%RECOV	%RECOV	%DIFF
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-001	BONCD1	Lead, Total	99.3	96.9	2.5

Recra LabNet - Lionville

INORGANICS PRECISION REPORT 09/29/98

CLIENT: TNU-HANFORD B98-063

RECRA LOT #: 9808L205

WORK ORDER: 10985-001-001-9999-00

SAMPLE	SITE ID	ANALYTE	INITIAL RESULT	REPLICATE	RPD	DILUTION FACTOR (REP)
-001REP	BONCD1	Lead, Total	4.8	2.4	66.7	1.0



Recra LabNet - Lionville

INORGANICS LABORATORY CONTROL STANDARDS REPORT 09/29/98

CLIENT: TNU-HANFORD B98-063

RECRA LOT #: 9808L205

WORK ORDER: 10985-001-001-9999-00

SAMPLE	SITE ID	ANALYTE	SAMPLE	SPIKED AMOUNT	SPIKED UNITS	%RECOV
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LCS1	98C0408-LC1	Mercury, LCS	5.4	5.0	UG/L	108.2
LCS1	98L1125-LC1	Lead, LCS	2490	2500	UG/L	99.7

Recra LabNet - Lionville Laboratory  
INORGANIC ANALYTICAL DATA PACKAGE FOR  
TNU-HANFORD B98-063

DATE RECEIVED: 08/05/98

RFW LOT # :9808L205

CLIENT ID /ANALYSIS	RFW #	MTX	PREP #	COLLECTION	EXTR/PREP	ANALYSIS
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BONCD1

MERCURY, TOTAL	001	W	98C0408	08/03/98	08/21/98	08/24/98
LEAD, TOTAL	001	W	98L1125	08/03/98	08/17/98	08/20/98
LEAD, TOTAL	001 REP	W	98L1125	08/03/98	08/17/98	08/20/98
LEAD, TOTAL	001 MS	W	98L1125	08/03/98	08/17/98	08/20/98
LEAD, TOTAL	001 MSD	W	98L1125	08/03/98	08/17/98	08/20/98

LAB QC:

MERCURY LABORATORY	LC1 BS	W	98C0408	N/A	08/21/98	08/24/98
MERCURY, TOTAL	MB1	W	98C0408	N/A	08/21/98	08/24/98
LEAD LABORATORY	LC1 BS	W	98L1125	N/A	08/17/98	08/20/98
LEAD, TOTAL	MB1	W	98L1125	N/A	08/17/98	08/20/98

**FIELD PERSONNEL: COMPLETE ONLY SHADED AREAS**

[illegible]

Bechtel Hanford Inc.		CHAIN OF CUSTODY/SAMPLE ANALYSIS REQUEST						B98-063-05		Page 1 of 1 <span style="float: right;">RUN 8/4/98</span>					
Collector Lahlberg/Nielson		Company Contact Steve Marske		Telephone No. 373-4316		Project Coordinator WEISS, RL		Data Turnaround  <b>21 Days</b>							
Project Designation 105-C Phase I & Phase II - Water Samples		Sampling Location 105-C		SAF No. B98-063											
Ice Chest No.		Field Logbook No. EL-1309-1		Method of Shipment Federal Express											
Shipped To <i>RUN REIRA LabNet</i> <i>Quanterra Incorporated</i> <i>2/3/98</i>		Offsite Property No.		Bill of Lading/Air Bill No.											
Waste Designation				COA											
POSSIBLE SAMPLE HAZARDS/REMARKS  <div style="text-align: center;">4235 7951 5429 <i>5.7</i></div>				Preservation		None	HNO3 to pH <2	Cool 4C	HNO3 to pH <2	HNO3 to pH <2	HNO3 to pH <2	Cool 4C	HNO3 to pH <2	HNO3 to pH <2	HNO3 to pH <2
				Type of Container		P	aG	P	P	P	P	aG	P	P	P
				No. of Container(s)		1	1	1	1	1	1	2	2	2	3
				Special Handling and/or Storage Cool 4C		Volume	20ml	500ml	500ml	500ml	1000ml	1000ml	1000ml	1000ml	1000ml
SAMPLE ANALYSIS				Activity Scan	Mercury - 7470 - (CV)	Chromium Hex - 7196	ICP Metals - 6010A (Supertrace) (1 cad)	See item (1) in Special Instructions	Nickel-63	PCBs - 8080 (Aroclor-1254)	See item (2) in Special Instructions	Isotopic Plutonium Isotopic Uranium Americium 241	Selenium-79		
Sample No.	Matrix *	Sample Date	Sample Time												
BONCD1	Water	8-3-98	1610		X	X	X			X					
CHAIN OF POSSESSION		Sign/Print Names				SPECIAL INSTRUCTIONS					Matrix *				
Relinquished By	Date/Time	Received By	Date/Time	(1) Gamma Spectroscopy (Cesium-137, Cobalt-60, Europium-152, Europium-154, Europium-155) (2) Gamma Spectroscopy (Water) (Cesium-137, Cobalt-60, Europium-152, Europium-154, Europium-155)  <i>Do not run Chromium-Hex 7196 unless directed by eec sample management.</i>					S Soil SE Sediment SO Solid SL Sludge W Water O Oil A Air DS Drum Solids DL Drum Liquid L Litter WI Waste L Liquid V Vegetation N Other						
<i>Steve Nielson</i>	<i>8/4/98 1300</i>	<i>FEA EY</i>													
Relinquished By	Date/Time	Received By	Date/Time												
Relinquished By	Date/Time	Received By	Date/Time												
Relinquished By	Date/Time	Received By	Date/Time												
Relinquished By	Date/Time	Received By	Date/Time												
LABORATORY SECTION	Received By	Title						Date/Time							
FINAL SAMPLE DISPOSITION	Disposal Method	Disposed By						Date/Time							